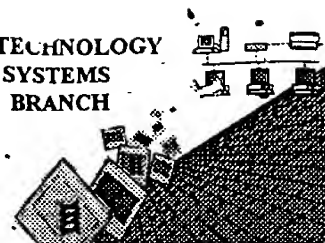


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BIOTECHNOLOGY  
SYSTEMS  
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**RAW SEQUENCE LISTING**  
**ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

09/831,426 B

Source:

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

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Revised 01/29/2002



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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/831,426B

DATE: 02/22/2002  
TIME: 15:16:38

Input Set : A:\9823seq.txt  
Output Set: N:\CRF3\02222002\I831426B.raw

**Does Not Comply  
Corrected Diskette Needed**

4 <110> APPLICANT: Hoechst Marion Roussel  
5 Bordon-Pallier, F.  
6 Rocher, C.  
8 <120> TITLE OF INVENTION: Human htFIIIA gene and coded htFIIIA protein  
10 <130> FILE REFERENCE: 146.1364  
12 <140> CURRENT APPLICATION NUMBER: US 09/831,426B  
13 <141> CURRENT FILING DATE: 2001-05-08  
15 <160> NUMBER OF SEQ ID NOS: 10  
17 <170> SOFTWARE: PatentIn Vers. 2.0

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**ERRORED SEQUENCES**

TECH CENTER 1600/2900

19 <210> SEQ ID NO: 1  
20 <211> LENGTH: 1273  
21 <212> TYPE: DNA  
22 <213> ORGANISM: Human  
24 <220> FEATURE:  
25 <221> NAME/KEY: CDS  
26 <222> LOCATION: (176)..(1270)  
E--> 28 <400> SEQUENCE: ①

*number for numeric identifier  
<400> should match numeric  
identifier <210>.*

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31 gtgcggcgct cgcgcgaagg ttcagcaggg agccgtgggc cgggcgcgcc ggttccgggc  
33 acgtgtctcg gcacgtggca gcgcgcctgg ccttgggctt ggaggcgccg gcgcc ctg  
34 Met  
35 1  
37 gat ccg ccg gcc gtg gtc gcc gag tgg gtg tgg tcc ttg acc atc gcc 226  
38 Asp Pro Pro Ala Val Val Ala Glu Ser Val Ser Ser Leu Thr Ile Ala  
39 5 10 15  
41 gac gcg ttc att gca gcc ggc gag agc tca gct ccg acc ccg ccg cgc 274  
42 Asp Ala Phe Ile Ala Ala Gly Glu Ser Ser Ala Pro Thr Pro Pro Arg  
43 20 25 30  
45 ccc gcg ctt ccc agg agg ttc atc tgc tcc ttc cct gac tgc agc gcc 322  
46 Pro Ala Leu Pro Arg Arg Phe Ile Cys Ser Phe Pro Asp Cys Ser Ala  
47 35 40 45  
49 aat tac agc aaa gcc tgg aag ctt gac gcg cac ctg tgc aag cac acg 370  
50 Asn Tyr Ser Lys Ala Trp Lys Leu Asp Ala His Leu Cys Lys His Thr  
51 50 55 60 65  
53 ggg gag aga cca ttt gtt tgt gac tat gaa ggg tgt ggc aag gcc ttc 418  
54 Gly Glu Arg Pro Phe Val Cys Asp Tyr Glu Gly Cys Gly Lys Ala Phe  
55 70 75 80  
57 atc agg gac tac cat ctg agc cgc cac att ctg act cac aca gga gaa 466  
58 Ile Arg Asp Tyr His Leu Ser Arg His Ile Leu Thr His Thr Gly Glu

*The type of errors shown exist throughout  
the Sequence Listing. Please check subse-  
quences for similar errors.*

## RAW SEQUENCE LISTING

DATE: 02/22/2002

PATENT APPLICATION: US/09/831,426B

TIME: 15:16:38

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Output Set: N:\CRF3\02222002\I831426B.raw

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62	Lys Pro Phe Val Cys Ala Ala Thr Gly Cys Asp Gln Lys Phe Asn Thr						
63	100 105 110						
65	aaa tca aac ttg aag aaa cat ttt gaa cgc aaa cat gaa aat caa caa	562					
66	Lys Ser Asn Leu Lys Lys His Phe Glu Arg Lys His Glu Asn Gln Gln						
67	115 120 125						
69	aaa caa tat ata tgc agt ttt gaa gac tgt aag aag acc ttt aag aaa	610					
70	Lys Gln Tyr Ile Cys Ser Phe Glu Asp Cys Lys Lys Thr Phe Lys Lys						
71	130 135 140 145						
73	cat cag cag ctg aaa atc cat cag tgc cag cat acc aat gaa cct cta	658					
74	His Gln Gln Leu Lys Ile His Gln Cys Gln His Thr Asn Glu Pro Leu						
75	150 155 160						
77	ttc aag tgt acc cag gaa gga tgt ggg aaa cac ttt gca tca ccc agc	706					
78	Phe Lys Cys Thr Gln Glu Gly Cys Gly Lys His Phe Ala Ser Pro Ser						
79	165 170 175						
81	aag ctg aaa cga cat gcc aag gcc cac gag ggc tat gta tgt caa aaa	754					
82	Lys Leu Lys Arg His Ala Lys Ala His Glu Gly Tyr Val Cys Gln Lys						
83	180 185 190						
85	gga tgt tcc ttt gtg gca aaa aca tgg acg gaa ctt ctg aaa cat gtg	802					
86	Gly Cys Ser Phe Val Ala Lys Thr Trp Thr Glu Leu Leu Lys His Val						
87	195 200 205						
89	aga gaa acc cat aaa gag gaa ata cta tgt gaa gta tgc cgg aaa aca	850					
90	Arg Glu Thr His Lys Glu Glu Ile Leu Cys Glu Val Cys Arg Lys Thr						
91	210 215 220 225						
93	ttt aaa cgc aaa gat tac ctt aag caa cac atg aaa act cat gcc cca	898					
94	Phe Lys Arg Lys Asp Tyr Leu Lys Gln His Met Lys Thr His Ala Pro						
95	230 235 240						
97	gaa agg gat gta tgt cgc tgt cca aga gaa ggc tgt gga aga acc tat	946					
98	Glu Arg Asp Val Cys Arg Cys Pro Arg Glu Gly Cys Gly Arg Thr Tyr						
99	245 250 255						
101	act act gtg ttt aat ctc caa agc cat atc ctc tcc ttc cat gag gaa	994					
102	Thr Thr Val Phe Asn Leu Gln Ser His Ile Leu Ser Phe His Glu Glu						
103	260 265 270						
105	agc cgc cct ttt gtg tgt gaa cat gct ggc tgt ggc aaa aca ttt gca	1042					
106	Ser Arg Pro Phe Val Cys Glu His Ala Gly Cys Gly Lys Thr Phe Ala						
107	275 280 285						
109	atg aaa caa agt ctc act agg cat gct gtt gta cat gat cct gac aag	1090					
110	Met Lys Gln Ser Leu Thr Arg His Ala Val Val His Asp Pro Asp Lys						
111	290 295 300 305						
113	aag aaa atg aag ctc aaa gtc aaa aaa tct cgt gaa aaa cgg agt ttg	1138					
114	Lys Lys Met Lys Leu Lys Val Lys Lys Ser Arg Glu Lys Arg Ser Leu						
115	310 315 320						
117	gcc tct cat ctc agt gga tat atc cct ccc aaa agg aaa caa ggg caa	1186					
118	Ala Ser His Leu Ser Gly Tyr Ile Pro Pro Lys Arg Lys Gln Gly Gln						
119	325 330 335						
121	ggc tta tct ttg tgt caa aac gga gag tca ccc aac tgt gtg gaa gac	1234					
122	Gly Leu Ser Leu Cys Gln Asn Gly Glu Ser Pro Asn Cys Val Glu Asp						
123	340 345 350						

## RAW SEQUENCE LISTING

DATE: 02/22/2002

PATENT APPLICATION: US/09/831,426B

TIME: 15:16:38

Input Set : A:\9823seq.txt

Output Set: N:\CRF3\02222002\I831426B.raw

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 131 <211> LENGTH: 365  
 132 <212> TYPE: PRT  
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 142 Arg Pro Ala Leu Pro Arg Arg Phe Ile Cys Ser Phe Pro Asp Cys Ser  
 143 35 40 45  
 145 Ala Asn Tyr Ser Lys Ala Trp Lys Leu Asp Ala His Leu Cys Lys His  
 146 50 55 60  
 148 Thr Gly Glu Arg Pro Phe Val Cys Asp Tyr Glu Gly Cys Gly Lys Ala  
 149 65 70 75 80  
 151 Phe Ile Arg Asp Tyr His Leu Ser Arg His Ile Leu Thr His Thr Gly  
 152 85 90 95  
 154 Glu Lys Pro Phe Val Cys Ala Ala Thr Gly Cys Asp Gln Lys Phe Asn  
 155 100 105 110  
 157 Thr Lys Ser Asn Leu Lys Lys His Phe Glu Arg Lys His Glu Asn Gln  
 158 115 120 125  
 160 Gln Lys Lys Tyr Ile Cys Ser Phe Glu Asp Cys Lys Lys Thr Phe Lys  
 161 130 135 140  
 163 Lys His Gln Gln Leu Lys Ile His Gln Cys Gln His Thr Asn Glu Pro  
 164 145 150 155 160  
 166 Leu Phe Lys Cys Thr Gln Glu Gly Cys Gly Lys His Phe Ala Ser Pro  
 167 165 170 175  
 169 Ser Lys Leu Lys Arg His Ala Lys Ala His Glu Gly Tyr Val Cys Gln  
 170 180 185 190  
 172 Lys Gly Cys Ser Phe Val Ala Lys Thr Trp Thr Glu Leu Leu Lys His  
 173 195 200 205  
 175 Val Arg Glu Thr His Lys Glu Ile Leu Cys Glu Val Cys Arg Lys  
 176 210 215 220  
 178 Thr Phe Lys Arg Lys Asp Tyr Leu Lys Gln His Met Lys Thr His Ala  
 179 225 230 235 240  
 181 Pro Glu Arg Asp Val Cys Arg Cys Pro Arg Glu Gly Cys Gly Arg Thr  
 182 245 250 255  
 184 Tyr Thr Thr Val Phe Asn Leu Gln Ser His Ile Leu Ser Phe His Glu  
 185 260 265 270  
 187 Glu Ser Arg Pro Phe Val Cys Glu His Ala Gly Cys Gly Lys Thr Phe  
 188 275 280 285  
 190 Ala Met Lys Gln Ser Leu Thr Arg His Ala Val Val His Asp Pro Asp  
 191 290 295 300  
 193 Lys Lys Lys Met Lys Leu Lys Val Lys Lys Ser Arg Glu Lys Arg Ser  
 194 305 310 315 320  
 196 Leu Ala Ser His Leu Ser Gly Tyr Ile Pro Pro Lys Arg Lys Gln Gly

DATE: 02/22/2002

TIME: 15:16:38

Input Set : A:\9823seq.txt

Output Set: N:\CRF3\02222002\I831426B.raw

197					325				330					335		
199	Gln	Gly	Leu	Ser	Leu	Cys	Gln	Asn	Gly	Glu	Ser	Pro	Asn	Cys	Val	Glu
200				340					345					350		
202	Asp	Lys	Met	Leu	Ser	Thr	Val	Ala	Val	Leu	Thr	Leu	Gly			
203			355					360					365			

206 <210> SEQ ID NO: 3

207 <211> LENGTH: 1273

208 <212> TYPE: DNA

209 <213> ORGANISM: Human

E--> 211 <400> SEQUENCE: 0

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214	gtgcggcgt	cgcggaagg	ttcagcagg	agcctgggc	cgggcgccc	ggttccggc	120
216	acgtgtctcg	gcacgtggca	gcgcgcctgg	ccctgggctt	ggaggcgccg	gcgccctgga	180
218	tccgcggcc	gtggctgcgc	agtcggtgtc	gtccttgacc	atcgccgacg	cgttcattgc	240
220	agcggcgag	agctcagctc	cgaccccgcc	gcgccccgcg	cttcccagga	ggttcattctg	300
222	ctccttcct	gactgcagcg	ccaattacag	caaagcctgg	aagcttgacg	cgcacctgtg	360
224	caagcacacg	ggggagagac	catttgtttg	tgactatgaa	gggtgtggca	aggccttcat	420
226	cagggaactac	catctgagcc	gccacattct	gactcacaca	ggagaaaagc	cgtttgtttg	480
228	tgcagccact	ggctgtgatc	aaaaattcaa	cacaaaaatca	aacttgaaga	aacattttga	540
230	acgcaaacat	gaaaatcaac	aaaaacaata	tatatgcagt	tttgaaact	gtaagaagac	600
232	ctttaaaaa	catcagcagc	tgaaaaatcca	tcagtgccag	cataccaatg	aacctctatt	660
234	caagtgtacc	caggaaggat	gtgggaaaca	ctttgcatca	cccagcaagc	tgaaacgaca	720
236	tgccaaggcc	cacgagggt	atgtatgtca	aaaaggatgt	tcctttgtgg	caaaaacatg	780
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240	ccggaaaaca	tttaaacgca	aagattacct	taagcaacac	atgaaaactc	atgcccaga	900
242	aagggatgta	tgctgctgtc	caagagaagg	ctgtggaaga	acctatacta	ctgtgtttaa	960
244	tctocaaagc	catatcctct	ccttccatga	ggaaagccgc	ccttttgtgt	gtgaacatgc	1020
246	tggctgtggc	aaaacatttg	caatgaaaca	aagtctcact	aggcatgctg	ttgtacatga	1080
248	tcctgacaag	aagaaaatga	agctcaaagt	caaaaaatct	cgtgaaaaac	ggagtttggc	1140
250	ctctcatctc	agtggatata	tccctcccaa	aaggaaacaa	gggcaaggct	tatctttgtg	1200
252	tcaaacatga	gagtcacca	actgtgtgga	agacaagatg	ctctcgacag	ttgcagtact	1260
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257 <210> SEO ID NO: 4

258 <211> LENGTH: 1213

259 <212> TYPE: DNA

260 <213> ORGANISM: ~~Human~~

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265	acgtgtctcg	gcacgtggca	gcgcgcctgg	ccctgggctt	ggaggcgccg	gcgccctgga	120
267	tccgcgggcc	gtggtcgcgc	agtcgggtgtc	gtccttgacc	atcgccgacg	cgttcattgc	180
269	agcgggcgag	agctcagctc	cgaccccgcc	gcgccccgcg	cttcccagga	ggttcacatc	240
271	ctccttccct	gactgcagcg	ccaattacag	caaagcctgg	aagcttgacg	cgcacctgtg	300
273	caagcacacg	gggagagagc	catttgtttg	tgactatgaa	gggtgtggca	aggccttcat	360
275	cagggaactac	catctgagcc	gccacattct	gactcacaca	ggagaaaagc	cgtttgtttg	420
277	tgcagccact	ggctgtgata	aaaaattcaa	cacaaaatca	aacttgaaga	aacattttga	480
279	acgcaaacat	gaaaatcaac	aaaaacaata	tatatgcagt	tttgaagact	gtaagaagac	540
281	ctttaagaaa	catcagcagc	tgaaaaatcca	tcagctgcag	cataccaatg	aacctctatt	600
283	caagtgtacc	caggaaggat	gtgggaaaca	ctttgcatac	cccagcaagc	tgaaacgaca	660
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## RAW SEQUENCE LISTING

DATE: 02/22/2002

PATENT APPLICATION: US/09/831,426B

TIME: 15:16:39

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291 aagggatgta tgcgctgtgc caagagaagg ctgtggaaga acctatacta ctgtgtttaa 900
293 tctccaaagc catatcctct ccttccatga ggaaagccgc ccttttgtgt gtgaacatgc 960
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299 ctctcatctc agtggatata tccctcccaa aaggaaacaa gggcaaggct tatctttgtg 1140
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325 <211> LENGTH: 20
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335 <212> TYPE: DNA
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357 cctcccgagg ccaagggtaa gtactgcaac 30
E--> 364 ①

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remove extra material at end of file.

## VERIFICATION SUMMARY

DATE: 02/22/2002

PATENT APPLICATION: US/09/831,426B

TIME: 15:16:40

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Output Set: N:\CRF3\02222002\I831426B.raw

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L:211 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:3 differs:0  
L:262 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:4 differs:0  
L:311 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:5 differs:0  
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L:347 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:9 differs:0  
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L:364 M:254 E: No. of Bases conflict, LENGTH:Input:7 Counted:30 SEQ:10